

SEQUENCE LISTING

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<120> METHODS AND COMPOSITIONS FOR MODULATING GAMETE ADHESION

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<160> 9

<170> PatentIn version 3.3

<210> 1
<211> 1281
<212> DNA
<213> Mus musculus

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180

aatgagactg agagaggacc atgctcccc aacccttgct acaatgatgc caaatgtctg
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gtgactttgg acacacagcg tggggacatc ttcaccgaat acatctgcca gtgccctgtg
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gccatttgctg attcacagat ttccgcctcg tctgtgtata tgggtttcat gggcttgcag
420

cgctggggcc cggagctggc tcgtctgtac cgcacagggta tcgtcaatgc ctggacagcc
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<210> 2
<211> 426
<212> PRT
<213> mus musculus

<400> 2

Met Gln Val Ser Arg Val Leu Ala Ala Leu Cys Gly Met Leu Leu Cys
1 5 10 15

Ala Ser Gly Leu Phe Ala Ala Ser Gly Asp Phe Cys Asp Ser Ser Leu
20 25 30

Cys Leu Asn Gly Gly Thr Cys Leu Thr Gly Gln Asp Asn Asp Ile Tyr
35 40 45

Cys Leu Cys Pro Glu Gly Phe Thr Gly Leu Val Cys Asn Glu Thr Glu
50 55 60

Arg Gly Pro Cys Ser Pro Asn Pro Cys Tyr Asn Asp Ala Lys Cys Leu
65 70 75 80

Val Thr Leu Asp Thr Gln Arg Gly Asp Ile Phe Thr Glu Tyr Ile Cys
85 90 95

Gln Cys Pro Val Gly Tyr Ser Gly Ile His Cys Glu Thr Gly Cys Ser

100

105

110

Thr Gln Leu Gly Met Glu Gly Gly Ala Ile Ala Asp Ser Gln Ile Ser
115 120 125

Ala Ser Ser Val Tyr Met Gly Phe Met Gly Leu Gln Arg Trp Gly Pro
130 135 140

Glu Leu Ala Arg Leu Tyr Arg Thr Gly Ile Val Asn Ala Trp Thr Ala
145 150 155 160

Ser Asn Tyr Asp Ser Lys Pro Trp Ile Gln Val Asn Leu Leu Arg Lys
165 170 175

Met Arg Val Ser Gly Val Met Thr Gln Gly Ala Ser Arg Ala Gly Arg
180 185 190

Ala Glu Tyr Leu Lys Thr Phe Lys Val Ala Tyr Ser Leu Asp Gly Arg
195 200 205

Lys Phe Glu Phe Ile Gln Asp Glu Ser Gly Gly Asp Lys Glu Phe Leu
210 215 220

Gly Asn Leu Asp Asn Asn Ser Leu Lys Val Asn Met Phe Asn Pro Thr
225 230 235 240

Leu Glu Ala Gln Tyr Ile Arg Leu Tyr Pro Val Ser Cys His Arg Gly
245 250 255

Cys Thr Leu Arg Phe Glu Leu Leu Gly Cys Glu Leu His Gly Cys Ser
260 265 270

Glu Pro Leu Gly Leu Lys Asn Asn Thr Ile Pro Asp Ser Gln Met Ser
275 280 285

Ala Ser Ser Ser Tyr Lys Thr Trp Asn Leu Arg Ala Phe Gly Trp Tyr
290 295 300

Pro His Leu Gly Arg Leu Asp Asn Gln Gly Lys Ile Asn Ala Trp Thr
305 310 315 320

Ala Gln Ser Asn Ser Ala Lys Glu Trp Leu Gln Val Asp Leu Gly Thr
325 330 335

Gln Arg Gln Val Thr Gly Ile Ile Thr Gln Gly Ala Arg Asp Phe Gly
340 345 350

His Ile Gln Tyr Val Ala Ser Tyr Lys Val Ala His Ser Asp Asp Gly
355 360 365

Val Gln Trp Thr Val Tyr Glu Glu Gln Gly Ser Ser Lys Val Phe Gln
370 375 380

Gly Asn Leu Asp Asn Asn Ser His Lys Lys Asn Ile Phe Glu Lys Pro
385 390 395 400

Phe Met Ala Arg Tyr Val Arg Val Leu Pro Val Ser Trp His Asn Arg
405 410 415

Ile Thr Leu Arg Leu Glu Leu Leu Gly Cys
420 425

<210> 3
<211> 404
<212> PRT
<213> mus musculus

<400> 3

Ala Ser Gly Asp Phe Cys Asp Ser Ser Leu Cys Leu Asn Gly Gly Thr
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Cys Leu Thr Gly Gln Asp Asn Asp Ile Tyr Cys Leu Cys Pro Glu Gly
20 25 30

Phe Thr Gly Leu Val Cys Asn Glu Thr Glu Arg Gly Pro Cys Ser Pro
35 40 45

Asn Pro Cys Tyr Asn Asp Ala Lys Cys Leu Val Thr Leu Asp Thr Gln
50 55 60

Arg Gly Asp Ile Phe Thr Glu Tyr Ile Cys Gln Cys Pro Val Gly Tyr
65 70 75 80

Ser Gly Ile His Cys Glu Thr Gly Cys Ser Thr Gln Leu Gly Met Glu
85 90 95

Gly Gly Ala Ile Ala Asp Ser Gln Ile Ser Ala Ser Ser Val Tyr Met
100 105 110

Gly Phe Met Gly Leu Gln Arg Trp Gly Pro Glu Leu Ala Arg Leu Tyr
115 120 125

Arg Thr Gly Ile Val Asn Ala Trp Thr Ala Ser Asn Tyr Asp Ser Lys
130 135 140

Pro Trp Ile Gln Val Asn Leu Leu Arg Lys Met Arg Val Ser Gly Val
145 150 155 160

Met Thr Gln Gly Ala Ser Arg Ala Gly Arg Ala Glu Tyr Leu Lys Thr
165 170 175

Phe Lys Val Ala Tyr Ser Leu Asp Gly Arg Lys Phe Glu Phe Ile Gln
180 185 190

Asp Glu Ser Gly Gly Asp Lys Glu Phe Leu Gly Asn Leu Asp Asn Asn
195 200 205

Ser Leu Lys Val Asn Met Phe Asn Pro Thr Leu Glu Ala Gln Tyr Ile
210 215 220

Arg Leu Tyr Pro Val Ser Cys His Arg Gly Cys Thr Leu Arg Phe Glu
225 230 235 240

Leu Leu Gly Cys Glu Leu His Gly Cys Ser Glu Pro Leu Gly Leu Lys
245 250 255

Asn Asn Thr Ile Pro Asp Ser Gln Met Ser Ala Ser Ser Ser Tyr Lys
260 265 270

Thr Trp Asn Leu Arg Ala Phe Gly Trp Tyr Pro His Leu Gly Arg Leu
275 280 285

Asp Asn Gln Gly Lys Ile Asn Ala Trp Thr Ala Gln Ser Asn Ser Ala
290 295 300

Lys Glu Trp Leu Gln Val Asp Leu Gly Thr Gln Arg Gln Val Thr Gly
305 310 315 320

Ile Ile Thr Gln Gly Ala Arg Asp Phe Gly His Ile Gln Tyr Val Ala
325 330 335

Ser Tyr Lys Val Ala His Ser Asp Asp Gly Val Gln Trp Thr Val Tyr
340 345 350

Glu Glu Gln Gly Ser Ser Lys Val Phe Gln Gly Asn Leu Asp Asn Asn
355 360 365

Ser His Lys Lys Asn Ile Phe Glu Lys Pro Phe Met Ala Arg Tyr Val

370

375

380

Arg Val Leu Pro Val Ser Trp His Asn Arg Ile Thr Leu Arg Leu Glu
385 390 395 400

Leu Leu Gly Cys

<210> 4
<211> 244
<212> PRT
<213> artificial

<220>
<223> EEC - recombinant protein

<400> 4

Ala Ser Gly Asp Phe Cys Asp Ser Ser Leu Cys Leu Asn Gly Gly Thr
1 5 10 15

Cys Leu Thr Gly Gln Asp Asn Asp Ile Tyr Cys Leu Cys Pro Glu Gly
20 25 30

Phe Thr Gly Leu Val Cys Asn Glu Thr Glu Arg Gly Pro Cys Ser Pro
35 40 45

Asn Pro Cys Tyr Asn Asp Ala Lys Cys Leu Val Thr Leu Asp Thr Gln
50 55 60

Arg Gly Asp Ile Phe Thr Glu Tyr Ile Cys Gln Cys Pro Val Gly Tyr
65 70 75 80

Ser Gly Ile His Cys Glu Thr Gly Cys Ser Thr Gln Leu Gly Met Glu
85 90 95

Gly Gly Ala Ile Ala Asp Ser Gln Ile Ser Ala Ser Ser Val Tyr Met
100 105 110

Gly Phe Met Gly Leu Gln Arg Trp Gly Pro Glu Leu Ala Arg Leu Tyr
115 120 125

Arg Thr Gly Ile Val Asn Ala Trp Thr Ala Ser Asn Tyr Asp Ser Lys
130 135 140

Pro Trp Ile Gln Val Asn Leu Leu Arg Lys Met Arg Val Ser Gly Val
145 150 155 160

Met Thr Gln Gly Ala Ser Arg Ala Gly Arg Ala Glu Tyr Leu Lys Thr
165 170 175

Phe Lys Val Ala Tyr Ser Leu Asp Gly Arg Lys Phe Glu Phe Ile Gln
180 185 190

Asp Glu Ser Gly Gly Asp Lys Glu Phe Leu Gly Asn Leu Asp Asn Asn
195 200 205

Ser Leu Lys Val Asn Met Phe Asn Pro Thr Leu Glu Ala Gln Tyr Ile
210 215 220

Arg Leu Tyr Pro Val Ser Cys His Arg Gly Cys Thr Leu Arg Phe Glu
225 230 235 240

Leu Leu Gly Cys

<210> 5
<211> 365
<212> PRT
<213> artificial

<220>
<223> ECC - recombinant protein

<400> 5

Glu Thr Glu Arg Gly Pro Cys Ser Pro Asn Pro Cys Tyr Asn Asp Ala
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Lys Cys Leu Val Thr Leu Asp Thr Gln Arg Gly Asp Ile Phe Thr Glu
20 25 30

Tyr Ile Cys Gln Cys Pro Val Gly Tyr Ser Gly Ile His Cys Glu Thr
35 40 45

Gly Cys Ser Thr Gln Leu Gly Met Glu Gly Gly Ala Ile Ala Asp Ser
50 55 60

Gln Ile Ser Ala Ser Ser Val Tyr Met Gly Phe Met Gly Leu Gln Arg
65 70 75 80

Trp Gly Pro Glu Leu Ala Arg Leu Tyr Arg Thr Gly Ile Val Asn Ala
85 90 95

Trp Thr Ala Ser Asn Tyr Asp Ser Lys Pro Trp Ile Gln Val Asn Leu
100 105 110

Leu Arg Lys Met Arg Val Ser Gly Val Met Thr Gln Gly Ala Ser Arg
115 120 125

Ala Gly Arg Ala Glu Tyr Leu Lys Thr Phe Lys Val Ala Tyr Ser Leu
130 135 140

Asp Gly Arg Lys Phe Glu Phe Ile Gln Asp Glu Ser Gly Gly Asp Lys
145 150 155 160

Glu Phe Leu Gly Asn Leu Asp Asn Asn Ser Leu Lys Val Asn Met Phe
165 170 175

Asn Pro Thr Leu Glu Ala Gln Tyr Ile Arg Leu Tyr Pro Val Ser Cys
180 185 190

His Arg Gly Cys Thr Leu Arg Phe Glu Leu Leu Gly Cys Glu Leu His
195 200 205

Gly Cys Ser Glu Pro Leu Gly Leu Lys Asn Asn Thr Ile Pro Asp Ser
210 215 220

Gln Met Ser Ala Ser Ser Ser Tyr Lys Thr Trp Asn Leu Arg Ala Phe
225 230 235 240

Gly Trp Tyr Pro His Leu Gly Arg Leu Asp Asn Gln Gly Lys Ile Asn
245 250 255

Ala Trp Thr Ala Gln Ser Asn Ser Ala Lys Glu Trp Leu Gln Val Asp
260 265 270

Leu Gly Thr Gln Arg Gln Val Thr Gly Ile Ile Thr Gln Gly Ala Arg
275 280 285

Asp Phe Gly His Ile Gln Tyr Val Ala Ser Tyr Lys Val Ala His Ser
290 295 300

Asp Asp Gly Val Gln Trp Thr Val Tyr Glu Glu Gln Gly Ser Ser Lys
305 310 315 320

Val Phe Gln Gly Asn Leu Asp Asn Asn Ser His Lys Lys Asn Ile Phe
325 330 335

Glu Lys Pro Phe Met Ala Arg Tyr Val Arg Val Leu Pro Val Ser Trp
340 345 350

His Asn Arg Ile Thr Leu Arg Leu Glu Leu Leu Gly Cys
355 360 365

<210> 6
<211> 205
<212> PRT
<213> artificial

<220>
<223> EC - recombinant protein

<400> 6

Glu Thr Glu Arg Gly Pro Cys Ser Pro Asn Pro Cys Tyr Asn Asp Ala
1 5 10 15

Lys Cys Leu Val Thr Leu Asp Thr Gln Arg Gly Asp Ile Phe Thr Glu
20 25 30

Tyr Ile Cys Gln Cys Pro Val Gly Tyr Ser Gly Ile His Cys Glu Thr
35 40 45

Gly Cys Ser Thr Gln Leu Gly Met Glu Gly Gly Ala Ile Ala Asp Ser
50 55 60

Gln Ile Ser Ala Ser Ser Val Tyr Met Gly Phe Met Gly Leu Gln Arg
65 70 75 80

Trp Gly Pro Glu Leu Ala Arg Leu Tyr Arg Thr Gly Ile Val Asn Ala
85 90 95

Trp Thr Ala Ser Asn Tyr Asp Ser Lys Pro Trp Ile Gln Val Asn Leu
100 105 110

Leu Arg Lys Met Arg Val Ser Gly Val Met Thr Gln Gly Ala Ser Arg
115 120 125

Ala Gly Arg Ala Glu Tyr Leu Lys Thr Phe Lys Val Ala Tyr Ser Leu
130 135 140

Asp Gly Arg Lys Phe Glu Phe Ile Gln Asp Glu Ser Gly Gly Asp Lys
145 150 155 160

Glu Phe Leu Gly Asn Leu Asp Asn Asn Ser Leu Lys Val Asn Met Phe
165 170 175

Asn Pro Thr Leu Glu Ala Gln Tyr Ile Arg Leu Tyr Pro Val Ser Cys
180 185 190

His Arg Gly Cys Thr Leu Arg Phe Glu Leu Leu Gly Cys
195 200 205

<210> 7
<211> 317
<212> PRT
<213> artificial

<220>
<223> CC - recombinant protein

<400> 7

Gly Cys Ser Thr Gln Leu Gly Met Glu Gly Gly Ala Ile Ala Asp Ser
1 5 10 15

Gln Ile Ser Ala Ser Ser Val Tyr Met Gly Phe Met Gly Leu Gln Arg
20 25 30

Trp Gly Pro Glu Leu Ala Arg Leu Tyr Arg Thr Gly Ile Val Asn Ala
35 40 45

Trp Thr Ala Ser Asn Tyr Asp Ser Lys Pro Trp Ile Gln Val Asn Leu
50 55 60

Leu Arg Lys Met Arg Val Ser Gly Val Met Thr Gln Gly Ala Ser Arg
65 70 75 80

Ala Gly Arg Ala Glu Tyr Leu Lys Thr Phe Lys Val Ala Tyr Ser Leu
85 90 95

Asp Gly Arg Lys Phe Glu Phe Ile Gln Asp Glu Ser Gly Gly Asp Lys
100 105 110

Glu Phe Leu Gly Asn Leu Asp Asn Asn Ser Leu Lys Val Asn Met Phe
115 120 125

Asn Pro Thr Leu Glu Ala Gln Tyr Ile Arg Leu Tyr Pro Val Ser Cys
130 135 140

His Arg Gly Cys Thr Leu Arg Phe Glu Leu Leu Gly Cys Glu Leu His
145 150 155 160

Gly Cys Ser Glu Pro Leu Gly Leu Lys Asn Asn Thr Ile Pro Asp Ser
165 170 175

Gln Met Ser Ala Ser Ser Ser Tyr Lys Thr Trp Asn Leu Arg Ala Phe
180 185 190

Gly Trp Tyr Pro His Leu Gly Arg Leu Asp Asn Gln Gly Lys Ile Asn
195 200 205

Ala Trp Thr Ala Gln Ser Asn Ser Ala Lys Glu Trp Leu Gln Val Asp
210 215 220

Leu Gly Thr Gln Arg Gln Val Thr Gly Ile Ile Thr Gln Gly Ala Arg
225 230 235 240

Asp Phe Gly His Ile Gln Tyr Val Ala Ser Tyr Lys Val Ala His Ser
245 250 255

Asp Asp Gly Val Gln Trp Thr Val Tyr Glu Glu Gln Gly Ser Ser Lys
260 265 270

Val Phe Gln Gly Asn Leu Asp Asn Asn Ser His Lys Lys Asn Ile Phe
275 280 285

Glu Lys Pro Phe Met Ala Arg Tyr Val Arg Val Leu Pro Val Ser Trp
290 295 300

His Asn Arg Ile Thr Leu Arg Leu Glu Leu Leu Gly Cys
305 310 315

<210> 8
<211> 25
<212> DNA
<213> artificial sequence

<220>
<223> primer sequence

<400> 8
cctcaggctg aggactggca gcggc

25

<210> 9
<211> 26
<212> DNA
<213> artificial sequence

<220>
<223> primer sequence

<400> 9
gctgtcaccg ggtgtccagg gtcacc

26